SEQUENCE LISTING

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<120> Agent and Method for Modulating Cell Migration

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<212> DNA
<213> Caenorhabditis elegans

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gct ctc ata ctc ctc gtc gtc tgc ctc gtt tat gcg ttg caa tca ggg    96
Ala Leu Ile Leu Leu Val Val Cys Leu Val Tyr Ala Leu Gln Ser Gly
             20                  25                  30

agt ggc acg atc tca gaa ttc tca tca gat gtg ctg ttc tcc agg gcc   144
Ser Gly Thr Ile Ser Glu Phe Ser Ser Asp Val Leu Phe Ser Arg Ala
         35                  40                  45

aag tac tca ggt gtg cca gtg cat cac agt cga tgg cgt caa gac gcc   192
Lys Tyr Ser Gly Val Pro Val His His Ser Arg Trp Arg Gln Asp Ala
     50                  55                  60

ggt ata cac gtc atc gac agc cat cac atc gtc cga aga gat tct tat   240
Gly Ile His Val Ile Asp Ser His His Ile Val Arg Arg Asp Ser Tyr
 65                  70                  75                  80

gga cgt cgt gga aaa cgt gat gtc acg tca aca gat cgg cga cgt cga   288
Gly Arg Arg Gly Lys Arg Asp Val Thr Ser Thr Asp Arg Arg Arg Arg
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ctc caa gga gtt gcc aga gac tgt gga cat gct tgt cac tta cga tta   336
Leu Gln Gly Val Ala Arg Asp Cys Gly His Ala Cys His Leu Arg Leu
            100                 105                 110
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     Arg Ser Asp Asp Ala Val Tyr Ile Val His Leu His Arg Trp Asn Gln
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     ata ccg gac tca cat aac aaa agt gtt ccc cac ttt tcc aat tca aat   432
  5  Ile Pro Asp Ser His Asn Lys Ser Val Pro His Phe Ser Asn Ser Asn
             130                 135                 140

     ttc gcg ccg atg gtc tta tat ttg gac tcg gag gag gag gtt aga ggt   480
     Phe Ala Pro Met Val Leu Tyr Leu Asp Ser Glu Glu Glu Val Arg Gly
     145                 150                 155                 160

 10  gga atg tct cga aca gat ccc gat tgt atc tac cgt gca cac gtt aaa   528
     Gly Met Ser Arg Thr Asp Pro Asp Cys Ile Tyr Arg Ala His Val Lys
                     165                 170                 175

     ggt gta cat cag cac agc atc gtc aat tta tgc gac tcg gaa gac gga   576
     Gly Val His Gln His Ser Ile Val Asn Leu Cys Asp Ser Glu Asp Gly
 15                  180                 185                 190

     ttg tac gga atg ctt gca cta ccc agc gga atc cat acg gtt gag cca   624
     Leu Tyr Gly Met Leu Ala Leu Pro Ser Gly Ile His Thr Val Glu Pro
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     att att agt gga aac gga aca gag cac gac gga gca agt cgc cat agg   672
 20  Ile Ile Ser Gly Asn Gly Thr Glu His Asp Gly Ala Ser Arg His Arg
             210                 215                 220

     caa cat ctc gtc cga aag ttc gat cca atg cac ttc aaa tcg ttt gac   720
     Gln His Leu Val Arg Lys Phe Asp Pro Met His Phe Lys Ser Phe Asp
     225                 230                 235                 240

 25  cat ctt aac tcg acc agt gtc aac gag acg gag acg acg gtt gcc acg   768
     His Leu Asn Ser Thr Ser Val Asn Glu Thr Glu Thr Thr Val Ala Thr
                     245                 250                 255

     tgg caa gat cag tgg gaa gat gtt att gaa cgc aaa gca aga tcc cga   816
     Trp Gln Asp Gln Trp Glu Asp Val Ile Glu Arg Lys Ala Arg Ser Arg
 30                  260                 265                 270

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     Arg Ala Ala Asn Ser Trp Asp His Tyr Val Glu Val Leu Val Val Ala
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     gat aca aaa atg tac gaa tat cac gga aga tct ctt gaa gac tac gtt   912
 35  Asp Thr Lys Met Tyr Glu Tyr His Gly Arg Ser Leu Glu Asp Tyr Val
             290                 295                 300

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     Leu Thr Leu Phe Ser Thr Val Ala Ser Ile Tyr Arg His Gln Ser Leu
     305                 310                 315                 320

 40  cgt gca tct atc aat gtc gtt gtt gtc aag ttg atc gtt ttg aaa acg  1008
     Arg Ala Ser Ile Asn Val Val Val Val Lys Leu Ile Val Leu Lys Thr
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  5 gat ttc tgt aga tgg cag cag tat tac aat gat cca gat gat tcg agt    1104
    Asp Phe Cys Arg Trp Gln Gln Tyr Tyr Asn Asp Pro Asp Asp Ser Ser
                355             360             365

    gtc caa cat cat gac gtt gca atc ctt ttg acg cgt aaa gat att tgt    1152
    Val Gln His His Asp Val Ala Ile Leu Leu Thr Arg Lys Asp Ile Cys
                370             375             380

 10 cga tca caa gga aaa tgc gat aca ctt gga ctt gct gaa ctt gga aca    1200
    Arg Ser Gln Gly Lys Cys Asp Thr Leu Gly Leu Ala Glu Leu Gly Thr
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    Met Cys Asp Met Gln Lys Ser Cys Ala Ile Ile Glu Asp Asn Gly Leu
 15                 405             410             415

    agt gct gca ttc aca att gct cat gaa ttg ggt cat gtg ttt tcg att    1296
    Ser Ala Ala Phe Thr Ile Ala His Glu Leu Gly His Val Phe Ser Ile
                420             425             430

    cct cat gat gac gaa cga aaa tgc tct acc tac atg ccg gtt aat aag    1344
 20 Pro His Asp Asp Glu Arg Lys Cys Ser Thr Tyr Met Pro Val Asn Lys
                435             440             445

    aac aac ttc cac ata atg gca cca acg ttg gaa tat aac act cat cca    1392
    Asn Asn Phe His Ile Met Ala Pro Thr Leu Glu Tyr Asn Thr His Pro
                450             455             460

 25 tgg agt tgg tcg cca tgt tca gct gga atg ctc gaa cga ttc ctc gaa    1440
    Trp Ser Trp Ser Pro Cys Ser Ala Gly Met Leu Glu Arg Phe Leu Glu
    465             470             475             480

    aat aat cga ggt caa act caa tgt cta ttc gat cag ccg gtc gaa cgt    1488
    Asn Asn Arg Gly Gln Thr Gln Cys Leu Phe Asp Gln Pro Val Glu Arg
 30                 485             490             495

    cgt tac tac gag gat gtc ttt gta cgt gat gaa cca gga aag aaa tac    1536
    Arg Tyr Tyr Glu Asp Val Phe Val Arg Asp Glu Pro Gly Lys Lys Tyr
                500             505             510

    gat gct cat caa cag tgc aag ttt gta ttt gga cca gct tct gag ttg    1584
 35 Asp Ala His Gln Gln Cys Lys Phe Val Phe Gly Pro Ala Ser Glu Leu
                515             520             525

    tgc cct tat atg ccg aca tgc cgc cgt ctt tgg tgt gca aca ttc tac    1632
    Cys Pro Tyr Met Pro Thr Cys Arg Arg Leu Trp Cys Ala Thr Phe Tyr
                530             535             540

 40 gga agc cag atg ggc tgt cga act cag cat atg cca tgg gcc gac gga    1680
    Gly Ser Gln Met Gly Cys Arg Thr Gln His Met Pro Trp Ala Asp Gly
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  5  Val Arg Leu Ala Pro Glu Ser Leu Thr Lys Ile Asp Gly Gln Trp Gly
                 580                 585                 590

     gac tgg cga tca tgg gga gaa tgc agt cgt act tgt ggt ggt ggt gtt    1824
     Asp Trp Arg Ser Trp Gly Glu Cys Ser Arg Thr Cys Gly Gly Gly Val
                 595                 600                 605

 10  caa aaa gga tta aga gat tgt gac agc cca aaa cct cga aat ggt gga    1872
     Gln Lys Gly Leu Arg Asp Cys Asp Ser Pro Lys Pro Arg Asn Gly Gly
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     aag tac tgt gtt ggt caa cga gaa cgt tat cgg tca tgt aat aca caa    1920
     Lys Tyr Cys Val Gly Gln Arg Glu Arg Tyr Arg Ser Cys Asn Thr Gln
 15  625                 630                 635                 640

     gaa tgc cca tgg gat act caa cca tac cgt gaa gtt caa tgt tct gaa    1968
     Glu Cys Pro Trp Asp Thr Gln Pro Tyr Arg Glu Val Gln Cys Ser Glu
                     645                 650                 655

     ttc aac aat aaa gat att gga atc caa ggt gtc gct tca acg aat act    2016
 20  Phe Asn Asn Lys Asp Ile Gly Ile Gln Gly Val Ala Ser Thr Asn Thr
                 660                 665                 670

     cac tgg gtt cca aaa tat gcg aat gtt gca cca aat gaa cgt tgc aag    2064
     His Trp Val Pro Lys Tyr Ala Asn Val Ala Pro Asn Glu Arg Cys Lys
                 675                 680                 685

 25  ctg tat tgt cgg ctc agt gga tct gca gcg ttc tat ctg ctt cga gat    2112
     Leu Tyr Cys Arg Leu Ser Gly Ser Ala Ala Phe Tyr Leu Leu Arg Asp
                 690                 695                 700

     aaa gtt gtt gat gga aca cca tgt gat aga aat gga gac gat att tgt    2160
     Lys Val Val Asp Gly Thr Pro Cys Asp Arg Asn Gly Asp Asp Ile Cys
 30  705                 710                 715                 720

     gta gct gga gct tgt atg cca gca ggc tgt gat cat caa ctt cat tca    2208
     Val Ala Gly Ala Cys Met Pro Ala Gly Cys Asp His Gln Leu His Ser
                     725                 730                 735

     act ctc cga aga gac aaa tgt ggt gtt tgc ggt ggg gat gat tct tcc    2256
 35  Thr Leu Arg Arg Asp Lys Cys Gly Val Cys Gly Gly Asp Asp Ser Ser
                 740                 745                 750

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     Cys Lys Val Val Lys Gly Thr Phe Asn Glu Gln Gly Thr Phe Gly Tyr
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 40  aac gaa gta atg aag att cca gct ggt tct gca aat att gat atc cgg    2352
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                 770                 775                 780
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      cgt gcc gcc aat ggt gaa ttc cta ctt aac ggt cat ttc caa gta tca   2448
  5   Arg Ala Ala Asn Gly Glu Phe Leu Leu Asn Gly His Phe Gln Val Ser
                      805             810             815

      ctg gct cgc caa caa att gca ttc caa gac act gtt ctc gaa tat tct   2496
      Leu Ala Arg Gln Gln Ile Ala Phe Gln Asp Thr Val Leu Glu Tyr Ser
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 10   ggt tct gat gca att att gaa cgg ata aat gga act ggt ccg att aga   2544
      Gly Ser Asp Ala Ile Ile Glu Arg Ile Asn Gly Thr Gly Pro Ile Arg
                  835             840             845

      agt gac att tat gtt cat gtt ctt tct gtt ggt agt cat cca ccc gac   2592
      Ser Asp Ile Tyr Val His Val Leu Ser Val Gly Ser His Pro Pro Asp
 15           850             855             860

      atc tca tat gag tac atg act gcg gct gtt cca aat gct gta att cgg   2640
      Ile Ser Tyr Glu Tyr Met Thr Ala Ala Val Pro Asn Ala Val Ile Arg
      865             870             875             880

      cca ata tcc agt gca ttg tat ttg tgg aga gtt acg gat act tgg aca   2688
 20   Pro Ile Ser Ser Ala Leu Tyr Leu Trp Arg Val Thr Asp Thr Trp Thr
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      gaa tgt gat aga gcc tgt cgt gga cag caa tcg caa aaa tta atg tgt   2736
      Glu Cys Asp Arg Ala Cys Arg Gly Gln Gln Ser Gln Lys Leu Met Cys
                  900             905             910

 25   ctg gac atg tcg act cat cgt caa agt cat gat aga aat tgt caa aat   2784
      Leu Asp Met Ser Thr His Arg Gln Ser His Asp Arg Asn Cys Gln Asn
                  915             920             925

      gtt ctc aaa cca aaa caa gca aca cga atg tgc aat ata gat tgt tct   2832
      Val Leu Lys Pro Lys Gln Ala Thr Arg Met Cys Asn Ile Asp Cys Ser
 30           930             935             940

      aca aga tgg atc act gaa gat gtg tct agt tgt agt gcc aaa tgt gga   2880
      Thr Arg Trp Ile Thr Glu Asp Val Ser Ser Cys Ser Ala Lys Cys Gly
      945             950             955             960

      tct gga cag aaa cgt caa cga gtt tct tgc gta aaa atg gag ggt gat   2928
 35   Ser Gly Gln Lys Arg Gln Arg Val Ser Cys Val Lys Met Glu Gly Asp
                      965             970             975

      cgt caa act cca gca tcc gaa cat cta tgt gat cgt aat tca aaa cca   2976
      Arg Gln Thr Pro Ala Ser Glu His Leu Cys Asp Arg Asn Ser Lys Pro
                  980             985             990

 40   tcc gat att gcc agt tgt tac att gac tgc tct gga aga aaa tgg aac   3024
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    atg cat cgg aag tca tat tgc gtt gat gat tcg aat cgt cga gtt gat    3120
  5 Met His Arg Lys Ser Tyr Cys Val Asp Asp Ser Asn Arg Arg Val Asp
    1025            1030                1035                1040

    gag tca ttg tgc ggc aga gaa cag aaa gag gcg aca gaa cgg gaa tgt    3168
    Glu Ser Leu Cys Gly Arg Glu Gln Lys Glu Ala Thr Glu Arg Glu Cys
                    1045                1050                1055

 10 aac aga att cca tgt cca aga tgg gtt tat ggg cat tgg tca gag tgc    3216
    Asn Arg Ile Pro Cys Pro Arg Trp Val Tyr Gly His Trp Ser Glu Cys
                1060                1065                1070

    tct cga agt tgt gat ggt gga gtc aaa atg cgt cat gct caa tgt ttg    3264
    Ser Arg Ser Cys Asp Gly Gly Val Lys Met Arg His Ala Gln Cys Leu
 15         1075                1080                1085

    gat gca gcc gat cgg gaa aca cat aca tcc aga tgt ggt cca gca cag    3312
    Asp Ala Ala Asp Arg Glu Thr His Thr Ser Arg Cys Gly Pro Ala Gln
        1090                1095                1100

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 20 Thr Gln Glu His Cys Asn Glu His Ala Cys Thr Trp Trp Gln Phe Gly
    1105                1110                1115                1120

    gtc tgg tct gac tgc tca gct aag tgt gga gat ggt gta cag tat cga    3408
    Val Trp Ser Asp Cys Ser Ala Lys Cys Gly Asp Gly Val Gln Tyr Arg
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 25 gac gct aat tgt acc gat cgt cat aga tca gta cta ccg gaa cat cgt    3456
    Asp Ala Asn Cys Thr Asp Arg His Arg Ser Val Leu Pro Glu His Arg
                1140                1145                1150

    tgc ctt aaa atg gaa aag ata att aca aaa cca tgt cat aga gaa tca    3504
    Cys Leu Lys Met Glu Lys Ile Ile Thr Lys Pro Cys His Arg Glu Ser
 30         1155                1160                1165

    tgt cca aaa tat aaa ctt gga gaa tgg tct cag tgt agt gtt tct tgt    3552
    Cys Pro Lys Tyr Lys Leu Gly Glu Trp Ser Gln Cys Ser Val Ser Cys
        1170                1175                1180

    gag gat gga tgg tcg tca aga aga gtt tca tgt gtt tct gga aat gga    3600
 35 Glu Asp Gly Trp Ser Ser Arg Arg Val Ser Cys Val Ser Gly Asn Gly
    1185                1190                1195                1200

    act gaa gtc gat atg tca ctt tgt ggt act gca tct gat cgg cct gct    3648
    Thr Glu Val Asp Met Ser Leu Cys Gly Thr Ala Ser Asp Arg Pro Ala
                    1205                1210                1215

 40 tct cat cag aca tgt aat tta ggc act tgc cca ttt tgg aga aat act    3696
    Ser His Gln Thr Cys Asn Leu Gly Thr Cys Pro Phe Trp Arg Asn Thr
                1220                1225                1230
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     aca acc gaa tgc ata tac cgc gaa caa tct gtt gat gct tct ttt tgt    3792
  5  Thr Thr Glu Cys Ile Tyr Arg Glu Gln Ser Val Asp Ala Ser Phe Cys
          1250                1255                1260

     gga gat acc aaa atg cca gaa act agt caa act tgc cat ctt ctg cca    3840
     Gly Asp Thr Lys Met Pro Glu Thr Ser Gln Thr Cys His Leu Leu Pro
     1265                1270                1275                1280

 10  tgt aca tct tgg aaa cca agt cat tgg tcc cct tgc tca gtc act tgt    3888
     Cys Thr Ser Trp Lys Pro Ser His Trp Ser Pro Cys Ser Val Thr Cys
                    1285                1290                1295

     gga tca gga att cag act aga agt gtt tcg tgt act cgt gga tct gaa    3936
     Gly Ser Gly Ile Gln Thr Arg Ser Val Ser Cys Thr Arg Gly Ser Glu
 15            1300                1305                1310

     gga act att gtt gat gaa tat ttt tgt gat cga aat act cgt cca cgc    3984
     Gly Thr Ile Val Asp Glu Tyr Phe Cys Asp Arg Asn Thr Arg Pro Arg
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     cta aaa aag act tgt gaa aaa gat act tgt gat ggg ccc aga gta ctt    4032
 20  Leu Lys Lys Thr Cys Glu Lys Asp Thr Cys Asp Gly Pro Arg Val Leu
          1330                1335                1340

     caa aaa ctt caa gcc gac gta cca cca atc cga tgg gca acc gga cca    4080
     Gln Lys Leu Gln Ala Asp Val Pro Pro Ile Arg Trp Ala Thr Gly Pro
     1345                1350                1355                1360

 25  tgg aca gcc tgt tca gca act tgt ggt aat ggt act caa cgt cgt ctt    4128
     Trp Thr Ala Cys Ser Ala Thr Cys Gly Asn Gly Thr Gln Arg Arg Leu
                    1365                1370                1375

     ctc aag tgc cga gat cat gtt cgt gat ctt cct gat gag tat tgc aat    4176
     Leu Lys Cys Arg Asp His Val Arg Asp Leu Pro Asp Glu Tyr Cys Asn
 30            1380                1385                1390

     cat ttg gat aag gaa gta tca aca aga aat tgt cgc ctt cgt gat tgt    4224
     His Leu Asp Lys Glu Val Ser Thr Arg Asn Cys Arg Leu Arg Asp Cys
          1395                1400                1405

     tca tac tgg aaa atg gcg gaa tgg gaa gag tgt cca gct act tgt gga    4272
 35  Ser Tyr Trp Lys Met Ala Glu Trp Glu Glu Cys Pro Ala Thr Cys Gly
          1410                1415                1420

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     Thr His Val Gln Gln Ser Arg Asn Val Thr Cys Val Ser Ala Glu Asp
     1425                1430                1435                1440

 40  ggt ggt cgg acg att ttg aaa gat gtt gat tgt gat gtg caa aag aga    4368
     Gly Gly Arg Thr Ile Leu Lys Asp Val Asp Cys Asp Val Gln Lys Arg
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    gaa cat att gga tcc tgg att att gga gat tgg tca aaa tgc tct gct    4464
  5 Glu His Ile Gly Ser Trp Ile Ile Gly Asp Trp Ser Lys Cys Ser Ala
            1475                 1480                 1485


    tct tgt ggt ggg gga tgg cgt cgt cgc agt gta tct tgc act tcg tct    4512
    Ser Cys Gly Gly Gly Trp Arg Arg Arg Ser Val Ser Cys Thr Ser Ser
            1490                 1495                 1500


 10 tct tgc gat gaa acc aga aaa cca aag atg ttt gat aaa tgc aat gaa    4560
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    1505             1510                 1515                 1520


    gaa cta tgt cca cca ctc aca aat aat tct tgg cag ata tct cca tgg    4608
    Glu Leu Cys Pro Pro Leu Thr Asn Asn Ser Trp Gln Ile Ser Pro Trp
 15                 1525                 1530                 1535


    act cac tgt tct gta tcg tgt ggc ggg gga gtt caa cgc cgc aaa atc    4656
    Thr His Cys Ser Val Ser Cys Gly Gly Gly Val Gln Arg Arg Lys Ile
                1540                 1545                 1550


    tgg tgt gaa gac gtg ctt tcc ggt cgt aaa caa gac gat atc gag tgc    4704
 20 Trp Cys Glu Asp Val Leu Ser Gly Arg Lys Gln Asp Asp Ile Glu Cys
            1555                 1560                 1565


    tca gag att aag cct cgc gaa caa aga gat tgt gaa atg cct cca tgc    4752
    Ser Glu Ile Lys Pro Arg Glu Gln Arg Asp Cys Glu Met Pro Pro Cys
            1570             1575                 1580


 25 cga tct cat tat cac aac aaa aca tca tca gca tca atg aca tca tta    4800
    Arg Ser His Tyr His Asn Lys Thr Ser Ser Ala Ser Met Thr Ser Leu
    1585             1590                 1595                 1600


    tca tct tcg aat tca aat acg acg tct tcc gct tcc gct tct tcg ctt    4848
    Ser Ser Ser Asn Ser Asn Thr Thr Ser Ser Ala Ser Ala Ser Ser Leu
 30                 1605                 1610                 1615


    cct atc ctt cca ccc gtc gtc tcc tgg caa acg tct gca tgg agc gcg    4896
    Pro Ile Leu Pro Pro Val Val Ser Trp Gln Thr Ser Ala Trp Ser Ala
                1620                 1625                 1630


    tgt tct gca aaa tgc ggt cgt gga acg aaa cga aga gtt gtc gaa tgt    4944
 35 Cys Ser Ala Lys Cys Gly Arg Gly Thr Lys Arg Arg Val Val Glu Cys
            1635                 1640                 1645


    gta aat cca tca tta aat gtg aca gtg gca agt aca gaa tgt gat caa    4992
    Val Asn Pro Ser Leu Asn Val Thr Val Ala Ser Thr Glu Cys Asp Gln
        1650                 1655                 1660


 40 acg aag aaa cca gtt gaa gaa gtt cgt tgt cgt act aaa cat tgc ccg    5040
    Thr Lys Lys Pro Val Glu Glu Val Arg Cys Arg Thr Lys His Cys Pro
    1665             1670                 1675                 1680


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     Arg Trp Lys Thr Thr Thr Trp Ser Ser Cys Ser Val Thr Cys Gly Arg
                 1685                1690                1695


     gga atc aga cgt cgt gaa gtt caa tgt tat cgt ggt cgc aag aat ttg        5136
  5  Gly Ile Arg Arg Arg Glu Val Gln Cys Tyr Arg Gly Arg Lys Asn Leu
                 1700                1705                1710


     gtg tct gat tcg gag tgc aat cca aaa act aag ctc aac tct gtt gcc        5184
     Val Ser Asp Ser Glu Cys Asn Pro Lys Thr Lys Leu Asn Ser Val Ala
                 1715                1720                1725


 10  aac tgt ttc cca gtg gct tgt cca gct tat aga tgg aat gtt act cca        5232
     Asn Cys Phe Pro Val Ala Cys Pro Ala Tyr Arg Trp Asn Val Thr Pro
              1730                1735                1740


     tgg agc aag tgc aaa gat gag tgt gct cga gga caa aag caa act cgt        5280
     Trp Ser Lys Cys Lys Asp Glu Cys Ala Arg Gly Gln Lys Gln Thr Arg
 15  1745                1750                1755                1760


     cgg gtg cac tgt ata agc act tct ggt aaa cga gca gct cca cga atg        5328
     Arg Val His Cys Ile Ser Thr Ser Gly Lys Arg Ala Ala Pro Arg Met
                 1765                1770                1775


     tgt gaa ttg gct cgt gca cca act tcg atc aga gag tgc gat aca tca        5376
 20  Cys Glu Leu Ala Arg Ala Pro Thr Ser Ile Arg Glu Cys Asp Thr Ser
                 1780                1785                1790


     aat tgt cca tat gag tgg gtg cca gga gat tgg caa acg tgt tca aag        5424
     Asn Cys Pro Tyr Glu Trp Val Pro Gly Asp Trp Gln Thr Cys Ser Lys
              1795                1800                1805


 25  tca tgt gga gaa gga gta cag aca cga gaa gtc aga tgt cgt aga aag        5472
     Ser Cys Gly Glu Gly Val Gln Thr Arg Glu Val Arg Cys Arg Arg Lys
              1810                1815                1820


     att aat ttt aac tca acc att cca att ata ttt atg ctc gaa gat gaa        5520
     Ile Asn Phe Asn Ser Thr Ile Pro Ile Ile Phe Met Leu Glu Asp Glu
 30  1825                1830                1835                1840


     cca gct gta cca aaa gag aaa tgt gaa ctt ttc cca aaa cca aat gaa        5568
     Pro Ala Val Pro Lys Glu Lys Cys Glu Leu Phe Pro Lys Pro Asn Glu
                 1845                1850                1855


     tct caa acg tgc gaa ctt aac cca tgc gat tcg gaa ttc aaa tgg agt        5616
 35  Ser Gln Thr Cys Glu Leu Asn Pro Cys Asp Ser Glu Phe Lys Trp Ser
                 1860                1865                1870


     ttc gga cca tgg ggt gaa tgc tcg aaa aat tgc ggt caa ggt att cga        5664
     Phe Gly Pro Trp Gly Glu Cys Ser Lys Asn Cys Gly Gln Gly Ile Arg
              1875                1880                1885


 40  cgt cga cgt gtc aag tgt gtg gcc aat gat ggt cgt cga gtt gaa cga        5712
     Arg Arg Arg Val Lys Cys Val Ala Asn Asp Gly Arg Arg Val Glu Arg
              1890                1895                1900
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      gtc aag tgt acc aca aag aaa cca cgt cga act caa tat tgt ttt gaa    5760
      Val Lys Cys Thr Thr Lys Lys Pro Arg Arg Thr Gln Tyr Cys Phe Glu
      1905                1910              1915                1920


      aga aat tgc ctt ccg tca act tgt cag gag ctt aaa tct cag aat gtt    5808
   5  Arg Asn Cys Leu Pro Ser Thr Cys Gln Glu Leu Lys Ser Gln Asn Val
                       1925              1930              1935


      aag gct aaa gat gga aat tac act att ctt ctt gac gga ttc act att    5856
      Lys Ala Lys Asp Gly Asn Tyr Thr Ile Leu Leu Asp Gly Phe Thr Ile
                  1940              1945              1950


  10  gaa att tat tgt cat cga atg aat tca acc att cct aaa gct tat ttg    5904
      Glu Ile Tyr Cys His Arg Met Asn Ser Thr Ile Pro Lys Ala Tyr Leu
                  1955              1960              1965


      aac gtt aat cca aga acc aat ttt gca gag gtt tat gga aaa aaa tta    5952
      Asn Val Asn Pro Arg Thr Asn Phe Ala Glu Val Tyr Gly Lys Lys Leu
  15       1970              1975              1980


      ata tac cct cat act tgc cca ttt aat ggt gat cgt aat gat tca tgc    6000
      Ile Tyr Pro His Thr Cys Pro Phe Asn Gly Asp Arg Asn Asp Ser Cys
      1985                1990              1995                2000


      cat tgt tca gaa gac ggc gat gca agt gct gga ttg acg aga ttc aat    6048
  20  His Cys Ser Glu Asp Gly Asp Ala Ser Ala Gly Leu Thr Arg Phe Asn
                       2005              2010              2015


      aaa gtt cga ata gat ttg ttg aat aga aag ttc cat ctg gcg gat tat    6096
      Lys Val Arg Ile Asp Leu Leu Asn Arg Lys Phe His Leu Ala Asp Tyr
                  2020              2025              2030


  25  aca ttt gca aaa cga gaa tat ggt gtt cat gtg cca tat ggt act gcc    6144
      Thr Phe Ala Lys Arg Glu Tyr Gly Val His Val Pro Tyr Gly Thr Ala
                  2035              2040              2045


      ggt gat tgc tac agt atg aaa gat tgt cca cag gga ata ttc tca att    6192
      Gly Asp Cys Tyr Ser Met Lys Asp Cys Pro Gln Gly Ile Phe Ser Ile
  30       2050              2055              2060


      gat tta aaa tct gct ggt ctg aaa tta gtt gac gat ctg aat tgg gag    6240
      Asp Leu Lys Ser Ala Gly Leu Lys Leu Val Asp Asp Leu Asn Trp Glu
      2065                2070              2075                2080


      gat caa ggt cat cga aca tcc tct cga atc gat cgt ttt tat aac aat    6288
  35  Asp Gln Gly His Arg Thr Ser Ser Arg Ile Asp Arg Phe Tyr Asn Asn
                       2085              2090              2095


      gca aaa gtt att ggt cac tgt ggt ggt ttt tgt gga aaa tgc tct cct    6336
      Ala Lys Val Ile Gly His Cys Gly Gly Phe Cys Gly Lys Cys Ser Pro
                  2100              2105              2110


  40  gag cgg tac aaa gga cta atc ttt gaa gtt aat aca aaa tta tta aat    6384
      Glu Arg Tyr Lys Gly Leu Ile Phe Glu Val Asn Thr Lys Leu Leu Asn
                  2115              2120              2125
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        cat gtg aaa aat ggt gga cac att gat gat gaa ttg gat gat gat ggt    6432
        His Val Lys Asn Gly Gly His Ile Asp Asp Glu Leu Asp Asp Asp Gly
            2130                2135                2140

        ttc tct ggt gac atg gat taatttttc gatacctaaa agtgtcaaaa            6480
   5    Phe Ser Gly Asp Met Asp
        2145                2150

        tctcgtatga atctctactt ctctggtctc ttatttcaag tttttgattc ttttcttttt  6540

        tttagttttt aatagcatta cttcgaattt attgtcattc cctcaatcac ctaacactag  6600

        gttttctaca tagtatgttc cttgaaaatg tttcatgatc aaaggttacg gtacttttg    6659


  10    <210> 2
        <211> 2150
        <212> PRT
        <213> Caenorhabditis elegans

        <400> 2
  15    Met Arg Ser Ile Gly Gly Ser Phe His Leu Leu Gln Pro Val Val Ala
          1               5                  10                  15

        Ala Leu Ile Leu Leu Val Val Cys Leu Val Tyr Ala Leu Gln Ser Gly
                        20                  25                  30

        Ser Gly Thr Ile Ser Glu Phe Ser Ser Asp Val Leu Phe Ser Arg Ala
  20              35                  40                  45

        Lys Tyr Ser Gly Val Pro Val His His Ser Arg Trp Arg Gln Asp Ala
                50                  55                  60

        Gly Ile His Val Ile Asp Ser His His Ile Val Arg Arg Asp Ser Tyr
           65                  70                  75                  80

  25    Gly Arg Arg Gly Lys Arg Asp Val Thr Ser Thr Asp Arg Arg Arg Arg
                        85                  90                  95

        Leu Gln Gly Val Ala Arg Asp Cys Gly His Ala Cys His Leu Arg Leu
                        100                 105                 110

        Arg Ser Asp Asp Ala Val Tyr Ile Val His Leu His Arg Trp Asn Gln
  30              115                 120                 125

        Ile Pro Asp Ser His Asn Lys Ser Val Pro His Phe Ser Asn Ser Asn
                130                 135                 140

        Phe Ala Pro Met Val Leu Tyr Leu Asp Ser Glu Glu Glu Val Arg Gly
        145                 150                 155                 160

  35    Gly Met Ser Arg Thr Asp Pro Asp Cys Ile Tyr Arg Ala His Val Lys
                        165                 170                 175

        Gly Val His Gln His Ser Ile Val Asn Leu Cys Asp Ser Glu Asp Gly
                        180                 185                 190
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      Leu Tyr Gly Met Leu Ala Leu Pro Ser Gly Ile His Thr Val Glu Pro
              195                 200                 205

      Ile Ile Ser Gly Asn Gly Thr Glu His Asp Gly Ala Ser Arg His Arg
              210                 215                 220

  5   Gln His Leu Val Arg Lys Phe Asp Pro Met His Phe Lys Ser Phe Asp
              225                 230                 235             240

      His Leu Asn Ser Thr Ser Val Asn Glu Thr Glu Thr Thr Val Ala Thr
                          245                 250                 255

 10   Trp Gln Asp Gln Trp Glu Asp Val Ile Glu Arg Lys Ala Arg Ser Arg
                      260                 265                 270

      Arg Ala Ala Asn Ser Trp Asp His Tyr Val Glu Val Leu Val Val Ala
                  275                 280                 285

      Asp Thr Lys Met Tyr Glu Tyr His Gly Arg Ser Leu Glu Asp Tyr Val
              290                 295                 300

 15   Leu Thr Leu Phe Ser Thr Val Ala Ser Ile Tyr Arg His Gln Ser Leu
              305                 310                 315             320

      Arg Ala Ser Ile Asn Val Val Val Val Lys Leu Ile Val Leu Lys Thr
                          325                 330                 335

      Glu Asn Ala Gly Pro Arg Ile Thr Gln Asn Ala Gln Gln Thr Leu Gln
 20                   340                 345                 350

      Asp Phe Cys Arg Trp Gln Gln Tyr Tyr Asn Asp Pro Asp Asp Ser Ser
                  355                 360                 365

      Val Gln His His Asp Val Ala Ile Leu Leu Thr Arg Lys Asp Ile Cys
              370                 375                 380

 25   Arg Ser Gln Gly Lys Cys Asp Thr Leu Gly Leu Ala Glu Leu Gly Thr
              385                 390                 395             400

      Met Cys Asp Met Gln Lys Ser Cys Ala Ile Ile Glu Asp Asn Gly Leu
                          405                 410                 415

      Ser Ala Ala Phe Thr Ile Ala His Glu Leu Gly His Val Phe Ser Ile
 30               420                 425                 430

      Pro His Asp Asp Glu Arg Lys Cys Ser Thr Tyr Met Pro Val Asn Lys
                  435                 440                 445

      Asn Asn Phe His Ile Met Ala Pro Thr Leu Glu Tyr Asn Thr His Pro
              450                 455                 460

 35   Trp Ser Trp Ser Pro Cys Ser Ala Gly Met Leu Glu Arg Phe Leu Glu
              465                 470                 475             480

      Asn Asn Arg Gly Gln Thr Gln Cys Leu Phe Asp Gln Pro Val Glu Arg
                      485                 490                 495
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      Arg Tyr Tyr Glu Asp Val Phe Val Arg Asp Glu Pro Gly Lys Lys Tyr
                  500                 505             510

      Asp Ala His Gln Gln Cys Lys Phe Val Phe Gly Pro Ala Ser Glu Leu
                  515             520             525

   5  Cys Pro Tyr Met Pro Thr Cys Arg Arg Leu Trp Cys Ala Thr Phe Tyr
                  530             535             540

      Gly Ser Gln Met Gly Cys Arg Thr Gln His Met Pro Trp Ala Asp Gly
      545             550             555             560

      Thr Pro Cys Asp Glu Ser Arg Ser Met Phe Cys His His Gly Ala Cys
  10                  565             570             575

      Val Arg Leu Ala Pro Glu Ser Leu Thr Lys Ile Asp Gly Gln Trp Gly
                  580             585             590

      Asp Trp Arg Ser Trp Gly Glu Cys Ser Arg Thr Cys Gly Gly Gly Val
                  595             600             605

  15  Gln Lys Gly Leu Arg Asp Cys Asp Ser Pro Lys Pro Arg Asn Gly Gly
                  610             615             620

      Lys Tyr Cys Val Gly Gln Arg Glu Arg Tyr Arg Ser Cys Asn Thr Gln
      625             630             635             640

      Glu Cys Pro Trp Asp Thr Gln Pro Tyr Arg Glu Val Gln Cys Ser Glu
  20                  645             650             655

      Phe Asn Asn Lys Asp Ile Gly Ile Gln Gly Val Ala Ser Thr Asn Thr
                  660             665             670

      His Trp Val Pro Lys Tyr Ala Asn Val Ala Pro Asn Glu Arg Cys Lys
                  675             680             685

  25  Leu Tyr Cys Arg Leu Ser Gly Ser Ala Ala Phe Tyr Leu Leu Arg Asp
                  690             695             700

      Lys Val Val Asp Gly Thr Pro Cys Asp Arg Asn Gly Asp Asp Ile Cys
      705             710             715             720

      Val Ala Gly Ala Cys Met Pro Ala Gly Cys Asp His Gln Leu His Ser
  30                  725             730             735

      Thr Leu Arg Arg Asp Lys Cys Gly Val Cys Gly Gly Asp Asp Ser Ser
                  740             745             750

      Cys Lys Val Val Lys Gly Thr Phe Asn Glu Gln Gly Thr Phe Gly Tyr
                  755             760             765

  35  Asn Glu Val Met Lys Ile Pro Ala Gly Ser Ala Asn Ile Asp Ile Arg
                  770             775             780

      Gln Lys Gly Tyr Asn Asn Met Lys Glu Asp Asp Asn Tyr Leu Ser Leu
      785             790             795             800
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        Arg Ala Ala Asn Gly Glu Phe Leu Leu Asn Gly His Phe Gln Val Ser
                        805                 810                 815

        Leu Ala Arg Gln Gln Ile Ala Phe Gln Asp Thr Val Leu Glu Tyr Ser
                        820                 825                 830

   5    Gly Ser Asp Ala Ile Ile Glu Arg Ile Asn Gly Thr Gly Pro Ile Arg
                        835                 840                 845

        Ser Asp Ile Tyr Val His Val Leu Ser Val Gly Ser His Pro Pro Asp
                    850                 855                 860

        Ile Ser Tyr Glu Tyr Met Thr Ala Ala Val Pro Asn Ala Val Ile Arg
  10    865                 870                 875                 880

        Pro Ile Ser Ser Ala Leu Tyr Leu Trp Arg Val Thr Asp Thr Trp Thr
                        885                 890                 895

        Glu Cys Asp Arg Ala Cys Arg Gly Gln Gln Ser Gln Lys Leu Met Cys
                    900                 905                 910

  15    Leu Asp Met Ser Thr His Arg Gln Ser His Asp Arg Asn Cys Gln Asn
                    915                 920                 925

        Val Leu Lys Pro Lys Gln Ala Thr Arg Met Cys Asn Ile Asp Cys Ser
                    930                 935                 940

        Thr Arg Trp Ile Thr Glu Asp Val Ser Ser Cys Ser Ala Lys Cys Gly
  20    945                 950                 955                 960

        Ser Gly Gln Lys Arg Gln Arg Val Ser Cys Val Lys Met Glu Gly Asp
                        965                 970                 975

        Arg Gln Thr Pro Ala Ser Glu His Leu Cys Asp Arg Asn Ser Lys Pro
                    980                 985                 990

  25    Ser Asp Ile Ala Ser Cys Tyr Ile Asp Cys Ser Gly Arg Lys Trp Asn
                    995                 1000                1005

        Tyr Gly Glu Trp Thr Ser Cys Ser Glu Thr Cys Gly Ser Asn Gly Lys
                1010                1015                1020

        Met His Arg Lys Ser Tyr Cys Val Asp Asp Ser Asn Arg Arg Val Asp
  30    1025                1030                1035                1040

        Glu Ser Leu Cys Gly Arg Glu Gln Lys Glu Ala Thr Glu Arg Glu Cys
                        1045                1050                1055

        Asn Arg Ile Pro Cys Pro Arg Trp Val Tyr Gly His Trp Ser Glu Cys
                    1060                1065                1070

  35    Ser Arg Ser Cys Asp Gly Gly Val Lys Met Arg His Ala Gln Cys Leu
                    1075                1080                1085

        Asp Ala Ala Asp Arg Glu Thr His Thr Ser Arg Cys Gly Pro Ala Gln
                1090                1095                1100
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Thr Gln Glu His Cys Asn Glu His Ala Cys Thr Trp Trp Gln Phe Gly
1105   1110   1115   1120

Val Trp Ser Asp Cys Ser Ala Lys Cys Gly Asp Gly Val Gln Tyr Arg
              1125   1130   1135

5 Asp Ala Asn Cys Thr Asp Arg His Arg Ser Val Leu Pro Glu His Arg
              1140   1145   1150

Cys Leu Lys Met Glu Lys Ile Ile Thr Lys Pro Cys His Arg Glu Ser
              1155   1160   1165

Cys Pro Lys Tyr Lys Leu Gly Glu Trp Ser Gln Cys Ser Val Ser Cys
10   1170   1175   1180

Glu Asp Gly Trp Ser Ser Arg Arg Val Ser Cys Val Ser Gly Asn Gly
1185   1190   1195   1200

Thr Glu Val Asp Met Ser Leu Cys Gly Thr Ala Ser Asp Arg Pro Ala
              1205   1210   1215

15 Ser His Gln Thr Cys Asn Leu Gly Thr Cys Pro Phe Trp Arg Asn Thr
              1220   1225   1230

Asp Trp Ser Ala Cys Ser Val Ser Cys Gly Ile Gly His Arg Glu Arg
              1235   1240   1245

Thr Thr Glu Cys Ile Tyr Arg Glu Gln Ser Val Asp Ala Ser Phe Cys
20   1250   1255   1260

Gly Asp Thr Lys Met Pro Glu Thr Ser Gln Thr Cys His Leu Leu Pro
1265   1270   1275   1280

Cys Thr Ser Trp Lys Pro Ser His Trp Ser Pro Cys Ser Val Thr Cys
              1285   1290   1295

25 Gly Ser Gly Ile Gln Thr Arg Ser Val Ser Cys Thr Arg Gly Ser Glu
              1300   1305   1310

Gly Thr Ile Val Asp Glu Tyr Phe Cys Asp Arg Asn Thr Arg Pro Arg
              1315   1320   1325

Leu Lys Lys Thr Cys Glu Lys Asp Thr Cys Asp Gly Pro Arg Val Leu
30   1330   1335   1340

Gln Lys Leu Gln Ala Asp Val Pro Pro Ile Arg Trp Ala Thr Gly Pro
1345   1350   1355   1360

Trp Thr Ala Cys Ser Ala Thr Cys Gly Asn Gly Thr Gln Arg Arg Leu
              1365   1370   1375

35 Leu Lys Cys Arg Asp His Val Arg Asp Leu Pro Asp Glu Tyr Cys Asn
              1380   1385   1390

His Leu Asp Lys Glu Val Ser Thr Arg Asn Cys Arg Leu Arg Asp Cys
              1395   1400   1405

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Ser Tyr Trp Lys Met Ala Glu Trp Glu Glu Cys Pro Ala Thr Cys Gly
  1410   1415   1420

Thr His Val Gln Gln Ser Arg Asn Val Thr Cys Val Ser Ala Glu Asp
  1425   1430   1435   1440

5 Gly Gly Arg Thr Ile Leu Lys Asp Val Asp Cys Asp Val Gln Lys Arg
                 1445   1450   1455

Pro Thr Ser Ala Arg Asn Cys Arg Leu Glu Pro Cys Pro Lys Gly Glu
             1460   1465   1470

Glu His Ile Gly Ser Trp Ile Ile Gly Asp Trp Ser Lys Cys Ser Ala
10   1475   1480   1485

Ser Cys Gly Gly Gly Trp Arg Arg Arg Ser Val Ser Cys Thr Ser Ser
        1490   1495   1500

Ser Cys Asp Glu Thr Arg Lys Pro Lys Met Phe Asp Lys Cys Asn Glu
  1505   1510   1515   1520

15 Glu Leu Cys Pro Pro Leu Thr Asn Asn Ser Trp Gln Ile Ser Pro Trp
                 1525   1530   1535

Thr His Cys Ser Val Ser Cys Gly Gly Gly Val Gln Arg Arg Lys Ile
             1540   1545   1550

Trp Cys Glu Asp Val Leu Ser Gly Arg Lys Gln Asp Asp Ile Glu Cys
20   1555   1560   1565

Ser Glu Ile Lys Pro Arg Glu Gln Arg Asp Cys Glu Met Pro Pro Cys
   1570   1575   1580

Arg Ser His Tyr His Asn Lys Thr Ser Ser Ala Ser Met Thr Ser Leu
  1585   1590   1595   1600

25 Ser Ser Ser Asn Ser Asn Thr Thr Ser Ser Ala Ser Ala Ser Ser Leu
                 1605   1610   1615

Pro Ile Leu Pro Pro Val Val Ser Trp Gln Thr Ser Ala Trp Ser Ala
             1620   1625   1630

Cys Ser Ala Lys Cys Gly Arg Gly Thr Lys Arg Arg Val Val Glu Cys
30   1635   1640   1645

Val Asn Pro Ser Leu Asn Val Thr Val Ala Ser Thr Glu Cys Asp Gln
   1650   1655   1660

Thr Lys Lys Pro Val Glu Glu Val Arg Cys Arg Thr Lys His Cys Pro
  1665   1670   1675   1680

35 Arg Trp Lys Thr Thr Thr Trp Ser Ser Cys Ser Val Thr Cys Gly Arg
                 1685   1690   1695

Gly Ile Arg Arg Arg Glu Val Gln Cys Tyr Arg Gly Arg Lys Asn Leu
             1700   1705   1710

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   Val Ser Asp Ser Glu Cys Asn Pro Lys Thr Lys Leu Asn Ser Val Ala
       1715                1720                1725

   Asn Cys Phe Pro Val Ala Cys Pro Ala Tyr Arg Trp Asn Val Thr Pro
       1730                1735                1740

 5 Trp Ser Lys Cys Lys Asp Glu Cys Ala Arg Gly Gln Lys Gln Thr Arg
       1745                1750                1755                1760

   Arg Val His Cys Ile Ser Thr Ser Gly Lys Arg Ala Ala Pro Arg Met
                   1765                1770                1775

   Cys Glu Leu Ala Arg Ala Pro Thr Ser Ile Arg Glu Cys Asp Thr Ser
10                 1780                1785                1790

   Asn Cys Pro Tyr Glu Trp Val Pro Gly Asp Trp Gln Thr Cys Ser Lys
       1795                1800                1805

   Ser Cys Gly Glu Gly Val Gln Thr Arg Glu Val Arg Cys Arg Arg Lys
       1810                1815                1820

15 Ile Asn Phe Asn Ser Thr Ile Pro Ile Ile Phe Met Leu Glu Asp Glu
       1825                1830                1835                1840

   Pro Ala Val Pro Lys Glu Lys Cys Glu Leu Phe Pro Lys Pro Asn Glu
                   1845                1850                1855

   Ser Gln Thr Cys Glu Leu Asn Pro Cys Asp Ser Glu Phe Lys Trp Ser
20                 1860                1865        .        1870

   Phe Gly Pro Trp Gly Glu Cys Ser Lys Asn Cys Gly Gln Gly Ile Arg
               1875                1880                1885

   Arg Arg Arg Val Lys Cys Val Ala Asn Asp Gly Arg Arg Val Glu Arg
           1890                1895                1900

25 Val Lys Cys Thr Thr Lys Lys Pro Arg Arg Thr Gln Tyr Cys Phe Glu
       1905                1910      .         1915                1920

   Arg Asn Cys Leu Pro Ser Thr Cys Gln Glu Leu Lys Ser Gln Asn Val
                   1925                1930                1935

   Lys Ala Lys Asp Gly Asn Tyr Thr Ile Leu Leu Asp Gly Phe Thr Ile
30                 1940                1945                1950

   Glu Ile Tyr Cys His Arg Met Asn Ser Thr Ile Pro Lys Ala Tyr Leu
       1955                1960                1965

   Asn Val Asn Pro Arg Thr Asn Phe Ala Glu Val Tyr Gly Lys Lys Leu
       1970                1975                1980

35 Ile Tyr Pro His Thr Cys Pro Phe Asn Gly Asp Arg Asn Asp Ser Cys
       1985                1990                1995                2000

   His Cys Ser Glu Asp Gly Asp Ala Ser Ala Gly Leu Thr Arg Phe Asn
                   2005                2010                2015
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Lys Val Arg Ile Asp Leu Leu Asn Arg Lys Phe His Leu Ala Asp Tyr
　　　　　　　2020　　　　　　　　　2025　　　　　　　2030

Thr Phe Ala Lys Arg Glu Tyr Gly Val His Val Pro Tyr Gly Thr Ala
　　　　　2035　　　　　　　　　2040　　　　　　　　2045

5　Gly Asp Cys Tyr Ser Met Lys Asp Cys Pro Gln Gly Ile Phe Ser Ile
　　　　　2050　　　　　　　2055　　　　　　　2060

Asp Leu Lys Ser Ala Gly Leu Lys Leu Val Asp Asp Leu Asn Trp Glu
　　2065　　　　　　　2070　　　　　　　　2075　　　　　　　2080

Asp Gln Gly His Arg Thr Ser Ser Arg Ile Asp Arg Phe Tyr Asn Asn
10　　　　　　　　2085　　　　　　　2090　　　　　　　2095

Ala Lys Val Ile Gly His Cys Gly Gly Phe Cys Gly Lys Cys Ser Pro
　　　　　　2100　　　　　　　2105　　　　　　　2110

Glu Arg Tyr Lys Gly Leu Ile Phe Glu Val Asn Thr Lys Leu Leu Asn
　　　　　2115　　　　　　　2120　　　　　　　2125

15　His Val Lys Asn Gly Gly His Ile Asp Asp Glu Leu Asp Asp Asp Gly
　　　　2130　　　　　　　2135　　　　　　　2140

Phe Ser Gly Asp Met Asp
2145　　　　　　2150